

SEQUENCE LISTING

<110> Myriad Genetics, Inc.
Cimbora, Daniel M.
Heichman, Karen
Bartel, Paul L.

<120> Protein-Protein Interactions

<130> 2318-282-II

<150> US 60/256,983

<151> 2000-12-21

<160> 4

<170> PatentIn version 3.0

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gaa Glu	aat Asn	ggg Gly 1070	gac Asp	gtc Val	ctc Leu	tta Leu	agc Ser 1075	ctg Leu	cag Gln	aga Arg	gct Ala	cat His 1080	gaa Glu	cag Gln	3789
gca Ala	gtg Val	aag Lys 1085	gaa Glu	aat Asn	gtg Val	aaa Lys	atg Met 1090	gct Ala	act Thr	gaa Glu	att Ile	tct Ser 1095	aga Arg	ttg Leu	3834
caa Gln	cag Gln	agg Arg 1100	cta Leu	caa Gln	aag Lys	tta Leu	gag Glu 1105	cca Pro	ggg Gly	tta Leu	gta Val	atg Met 1110	tct Ser	tct Ser	3879
tgt Cys	ttg Leu	gat Asp 1115	gag Glu	cca Pro	gct Ala	act Thr	gag Glu 1120	ttt Phe	ttt Phe	gga Gly	aat Asn	act Thr 1125	gcg Ala	gaa Glu	3924
caa Gln	aca Thr	gag Glu 1130	cag Gln	ttt Phe	tta Leu	cag Gln	caa Gln 1135	aac Asn	cga Arg	acg Thr	aag Lys	caa Gln 1140	gta Val	gaa Glu	3969
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gtc Val	cgg Arg	gac Asp 1160	ctg Leu	gga Gly	agt Ser	aca Thr	ggg Gly 1165	acg Thr	agc Ser	tct Ser	gtt Val	cag Gln 1170	aga Arg	cag Gln	4059
gaa Glu	gtc Val	aaa Lys 1175	ata Ile	gag Glu	gag Glu	tct Ser	gaa Glu 1180	gct Ala	tca Ser	gta Val	gag Glu	ggg Gly 1185	ttt Phe	tct Ser	4104
gag Glu	ctt Leu	gaa Glu 1190	aac Asn	agt Ser	gaa Glu	gag Glu	acc Thr 1195	agg Arg	act Thr	gaa Glu	tcc Ser	tgg Trp 1200	gag Glu	ctg Leu	4149
aag Lys	aat Asn	cag Gln 1205	att Ile	agt Ser	cag Gln	ctt Leu	cag Gln 1210	gaa Glu	cag Gln	cta Leu	atg Met	atg Met 1215	tta Leu	tgt Cys	4194
gcg Ala	gac Asp	tgt Cys 1220	gat Asp	cga Arg	gct Ala	tct Ser	gaa Glu 1225	aag Lys	aaa Lys	cag Gln	gac Asp	cta Leu 1230	ctt Leu	ttt Phe	4239
gat Asp	gtt Val	tct Ser 1235	gtg Val	cta Leu	aaa Lys	aag Lys	aaa Lys 1240	ctg Leu	aag Lys	atg Met	ctt Leu	gag Glu 1245	aga Arg	atc Ile	4284
cct Pro	gag Glu	gct Ala 1250	tct Ser	ccc Pro	aaa Lys	tat Tyr	aag Lys 1255	ctg Leu	ttg Leu	tat Tyr	gaa Glu	gat Asp 1260	gtg Val	agc Ser	4329

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Leu Arg Tyr Glu Ser Glu Lys Leu Gln Gln Glu Asn Ser Ile Leu	
1520 1525 1530	
aga aat gaa att act act tta aat gaa gaa gat agc att tct aac	5184
Arg Asn Glu Ile Thr Thr Leu Asn Glu Glu Asp Ser Ile Ser Asn	
1535 1540 1545	
ctg aaa tta ggg aca tta aat gga tct cag gaa gaa atg tgg caa	5229
Leu Lys Leu Gly Thr Leu Asn Gly Ser Gln Glu Glu Met Trp Gln	
1550 1555 1560	
aaa acg gaa act gta aaa caa gaa aat gct gca gtt cag aag atg	5274
Lys Thr Glu Thr Val Lys Gln Glu Asn Ala Ala Val Gln Lys Met	
1565 1570 1575	
gtt gaa aat tta aag aaa cag att tca gaa tta aaa atc aaa aac	5319
Val Glu Asn Leu Lys Lys Gln Ile Ser Glu Leu Lys Ile Lys Asn	
1580 1585 1590	
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Gln Gln Leu Asp Leu Glu Asn Thr Glu Leu Ser Gln Lys Asn Ser	
1595 1600 1605	
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Gln Asn Gln Glu Lys Leu Gln Glu Leu Asn Gln Arg Leu Thr Glu	
1610 1615 1620	
atg cta tgc cag aag gaa aaa gag cca gga aac agt gca ttg gag	5454
Met Leu Cys Gln Lys Glu Lys Glu Pro Gly Asn Ser Ala Leu Glu	
1625 1630 1635	
gaa cgg gaa caa gag aag ttt aat ctg aaa gaa gaa ctg gaa cgt	5499
Glu Arg Glu Gln Glu Lys Phe Asn Leu Lys Glu Glu Leu Glu Arg	
1640 1645 1650	
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Cys Lys Val Gln Ser Ser Thr Leu Val Ser Ser Leu Glu Ala Glu	
1655 1660 1665	
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Leu Ser Glu Val Lys Ile Gln Thr His Ile Val Gln Gln Glu Asn	
1670 1675 1680	
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His Leu Leu Lys Asp Glu Leu Glu Lys Met Lys Gln Leu His Arg	
1685 1690 1695	
tgt ccc gat ctc tct gac ttc cag caa aaa atc tct agt gtt cta	5679
Cys Pro Asp Leu Ser Asp Phe Gln Gln Lys Ile Ser Ser Val Leu	
1700 1705 1710	
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Ser Tyr Asn Glu Lys Leu Leu Lys Glu Lys Glu Ala Leu Ser Glu	
1715 1720 1725	
gaa tta aat agc tgt gtc gat aag ttg gca aaa tca agt ctt tta	5769
Glu Leu Asn Ser Cys Val Asp Lys Leu Ala Lys Ser Ser Leu Leu	
1730 1735 1740	

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Lys Val Gln Asn Leu Glu Asp Thr Val Gln Asn Val Asn Leu Gln	
1775 1780 1785	
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Met Ser Arg Met Lys Ser Asp Leu Arg Val Thr Gln Gln Glu Lys	
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Glu Ala Leu Lys Gln Glu Val Met Ser Leu His Lys Gln Leu Gln	
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Ser Gly Leu His Asn Gln Gln Lys Arg Leu Ser Trp Asp Lys Leu	
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Asp His Leu Met Asn Glu Glu Gln Gln Leu Leu Trp Gln Glu Asn	
1850 1855 1860	
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Glu Arg Leu Gln Thr Met Val Gln Asn Thr Lys Ala Glu Leu Thr	
1865 1870 1875	
cac tcc cgg gag aag gtc cgt caa ttg gaa tcc aat ctt ctt ccc	6219
His Ser Arg Glu Lys Val Arg Gln Leu Glu Ser Asn Leu Leu Pro	
1880 1885 1890	
aag cac caa aaa cat cta aac cca tca ggt acc atg aat ccc aca	6264
Lys His Gln Lys His Leu Asn Pro Ser Gly Thr Met Asn Pro Thr	
1895 1900 1905	
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Glu Gln Glu Lys Leu Ser Leu Lys Arg Glu Cys Asp Gln Phe Gln	
1910 1915 1920	
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Lys Glu Gln Ser Pro Ala Asn Arg Lys Val Ser Gln Met Asn Ser	
1925 1930 1935	
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Leu Glu Gln Glu Leu Glu Thr Ile His Leu Glu Asn Glu Gly Leu	
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Lys Lys Lys Gln Val Lys Leu Asp Glu Gln Leu Met Glu Met Gln	
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His Leu Arg Ser Thr Ala Thr Pro Ser Pro Ser Pro His Ala Trp	
1970 1975 1980	

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Arg Arg Ser Leu Pro Glu Phe Gln Glu Ser Val Glu Glu Phe Pro Glu

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